

JC20 Rec'd PCT/PTO

Attorney Docket No. Le A 32 70



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

re application of: Burchardt, et al.

Application Serial No.: 09/701,313

Filed: 05/18/2001

Title: Monoclonal Antibody and Assay for Detecting PliiNP

CERTIFICATION OF MAILING UNDER 37 C.F.R. 1.8(a)

I hereby certify that this correspondence and any papers referred to as attached are being deposited, on the date shown below, with the United States Postal Service, with sufficient postage, as first class mail in an envelope addressed to Box Sequence, Assistant Commissioner for Patents, Washington, D.C. 20231.

Date:

JAN 3 0 2002

Box Sequence Assistant Commissioner for Patents Washington, D.C. 20231

RESPONSE TO NOTIFICATION OF MISSING REQUIREMENTS UNDER 35 U.S.C. 371 IN THE UNITED STATES DESIGNATED/ELECTED OFFICE (DO/EO/US)

Sir:

This replies to the Notification of Missing Requirements mailed on 6 December 2001. A copy of the Notification is enclosed.

Submitted herewith is a:

- 1. Substitute copy of the Sequence Listing for the nucleotide and/or amino acid sequence(s) in this application.
- 2. Please insert the substitute copy of the sequence listing into the specification.
- 3. The computer readable form and paper copy submitted herewith are the same, and introduce no new matter over what is disclosed in the application as originally filed (37 C.F.R. 1.821(g)).

Respectfully submitted,

Reg. No.: 48,972

Tel. No.: (203) 812-6450

Date: JAN 3 0 2002

Susan M. Pellegrino

Attorney for Applicant **Bayer Corporation**

400 Morgan Lane

West Haven, CT 06516



Convessioner for Patents, Box PCT United States Valent and Tradamark Office Washington, IAC, 20231

ATTY. DOCKET NO. FIRST NAMED APPLICANT U.S. APPLICATION NUMBER NO. LeA 32 701 Elmar Reinhold Burchardt 09/701.313

INTERNATIONAL APPLICATION NO.

PCT/EP99/03392

I.A. FILING DATE

PRIORITY DATE

05/17/1999

05/28/1998

CONFIRMATION NO. 8752 371 FORMALITIES LETTER

OC000000007155420*

Jeffrey M Greenman Vice President Patents and Licensing **Bayer Corporation** 400 Morgan Lane West Haven, CT 06516

Date Mailed: 12/06/2001

NOTIFICATION OF MISSING REQUIREMENTS UNDER 35 U.S.C. 371 IN THE UNITED STATES DESIGNATED/ELECTED OFFICE (DO/EO/US)

The following items have been submitted by the applicant or the IB to the United States Patent and Trademark Office as an Elected Office (37 CFR 1.495):

- U.S. Basic National Fees
- Priority Document
- Biochemical Sequence Diskette
- Biochemical Sequence Listing
- Copy of IPE Report
- · Copy of references cited in ISR
- Copy of the International Application
- Copy of the International Search Report
- Information Disclosure Statements
- Oath or Declaration
- Request for Immediate Examination

RECEIVED

DEC 1 4 2001

JEFFREY M. GREENMAN

The following items MUST be furnished within the period set forth below in order to complete the requirements for acceptance under 35 U.S.C. 371:

- The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821-1.825 for the following reason(s):
 - A copy of the "Sequence Listing" in computer readable form has been submitted. The content of the computer readable form, however, does not comply with the requirements of 37 CFR 1.822 and/or 1.832, as indicated on the attached marked-up copy of the "Raw Sequence Listing."
 - APPLICANT MUST PROVIDE:
 - An initial or substitute paper copy or compact disc of the "Sequence Listing," as well as an amendment directing its entry into the specification.
 - A statement that the contents of the paper or compact disc and the computer readable form are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e),

1.821(f), 1.821(g), 1.825(b) or 1.825(d).

- For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:
 - For Rules Interpretation, call (703) 308-4216
 - To Purchase Patentin Software, call (703) 306-2600
 - For Patentin Software Program Help, call (703) 306-4119 or e-mail at patin21help@uspto.gov or patin3help@uspto.gov

ALL OF THE ITEMS SET FORTH ABOVE MUST BE SUBMITTED WITHIN TWO (2) MONTH FROM THE DATE OF THIS NOTICE OR BY 22 or 32 MONTHS (where 37 CFR 1.495 applies) FROM THE PRIORITY DATE FOR THE APPLICATION, WHICHEVER IS LATER. FAILURE TO PROPERLY RESPOND WILL RESULT IN ABANDONMENT.

The time period set above may be extended by filing a petition and fee for extension of time under the provisions of 37 CFR 1.136(a).

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

A copy of this notice MUST be returned with the response.

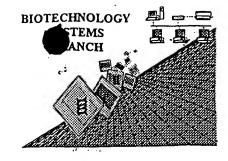
SHAKEEL AHMED

Telephone: (703) 305-3659

PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
09/701 313	PCT/EP99/03392	LeA 32 701

RAW SEQUENCE LISTING ERROR REPORT



SK

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/701, 3/3Source: Pcr/09Date Processed by STIC: 9/5/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,313

DATE: 07/05/2001 TIME: 15:51:29

Input Set : A:\32701.app

Output Set: N:\CRF3\07032001\I701313.raw

Does Not Comply Corrected Diskette Needed

3 <110> APPLICANT: Bayer AG

5 <120> TITLE OF INVENTION: Monoclonal antibody and assay for detecting PIIINP

7 <130> FILE REFERENCE: MoAb and assay for detecting PIIINP

9 <140 > CURRENT APPLICATION NUMBER: US/09/701,313

12 <160> NUMBER OF SEQ ID NOS: 13

14 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

189 <210> SEQ ID NO: 13 190 <211> LENGTH (31) 30 slown

191 <212> TYPE: DNA

192 <213> ORGANISM: 'Axial Seamount' polynoid polychaete

194 <400> SEQUENCE: 13

E--> 195 cgcgaagctt gggagaatag ttctgaggac

30

VERIFICATION SUMMARY PATENT APPLICATION: US/09/701,313

DATE: 07/05/2001 TIME: 15:51:30

Input Set : A:\32701.app

Output Set: N:\CRF3\07032001\1701313.raw

9 M:270 C: Current Application Number differs, Replaced Current Application Number 10 M:271 C: Current Filing Date differs, Replaced Current Filing Date 195 M:252 E: No. of Seq. differs, <211>LENGTH:Input:31 Found:30 SEQ:13